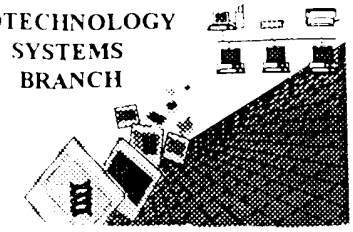
BIOTECHNOLOGY SYSTEMS BRANCH



217

1651

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/925,055B
Source: 1647
Date Processed by STIC: 2/19/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 3.1 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**http://www.uspto.gov/ebc/efs/downloads/documents.htm**> , **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. **Hand Carry directly to:**
   U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
   Or
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
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Revised 01/29/2002

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 09/925,055B

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1_____Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2_____Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3_____Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4_____Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5_____Variable Length	Sequence(s)____contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6_____PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7_____Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8_____Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9_____Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10_____Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
11 J Use of <220>	Sequence(s) 30 missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12_____PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13_____Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

1644

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                    RAW SEQUENCE LISTING                      DATE: 02/19/2002
                    PATENT APPLICATION:  US/09/925,055B        TIME: 12:20:18

                    Input Set : A:\00-56 SEQ.txt
                    Output Set: N:\CRF3\02192002\I925055B.raw

      3 <110> APPLICANT: Kindsvogel, Wayne R.
      4       Topouzis, Stavros
      6 <120> TITLE OF INVENTION: SOLUBLE ZCYTOR11 CYTOKINE RECEPTORS
      8 <130> FILE REFERENCE: 00-56
C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/925,055B
C--> 10 <141> CURRENT FILING DATE: 2001-08-08
     10 <150> PRIOR APPLICATION NUMBER: US 60/223,827
     11 <151> PRIOR FILING DATE: 2000-08-08
     13 <150> PRIOR APPLICATION NUMBER: US 60/250,876
     14 <151> PRIOR FILING DATE: 2000-12-01
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     18 <170> SOFTWARE: FastSEQ for Windows Version 3.0
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     22 <212> TYPE: DNA
     23 <213> ORGANISM: Homo sapien
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     27 <222> LOCATION: (34)...(1755)
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     31                                      Met Arg Thr Leu Leu Thr Ile
     32                                       1               5
     34 ttg act gtg gga tcc ctg gct gct cac gcc cct gag gac ccc tcg gat     102
     35 Leu Thr Val Gly Ser Leu Ala Ala His Ala Pro Glu Asp Pro Ser Asp
     36         10                  15                  20
     38 ctg ctc cag cac gtg aaa ttc cag tcc agc aac ttt gaa aac atc ctg     150
     39 Leu Leu Gln His Val Lys Phe Gln Ser Ser Asn Phe Glu Asn Ile Leu
     40     25                  30                  35
     42 acg tgg gac agc ggg cca gag ggc acc cca gac acg gtc tac agc atc     198
     43 Thr Trp Asp Ser Gly Pro Glu Gly Thr Pro Asp Thr Val Tyr Ser Ile
     44  40                  45                  50                  55
     46 gag tat aag acg tac gga gag agg gac tgg gtg gca aag aag ggc tgt     246
     47 Glu Tyr Lys Thr Tyr Gly Glu Arg Asp Trp Val Ala Lys Lys Gly Cys
     48                 60                  65                  70
     50 cag cgg atc acc cgg aag tcc tgc aac ctg acg gtg gag acg ggc aac     294
     51 Gln Arg Ile Thr Arg Lys Ser Cys Asn Leu Thr Val Glu Thr Gly Asn
     52             75                  80                  85
     54 ctc acg gag ctc tac tat gcc agg gtc acc gct gtc agt gcg gga ggc     342
     55 Leu Thr Glu Leu Tyr Tyr Ala Arg Val Thr Ala Val Ser Ala Gly Gly
     56         90                  95                 100
     58 cgg tca gcc acc aag atg act gac agg ttc agc tct ctg cag cac act     390
     59 Arg Ser Ala Thr Lys Met Thr Asp Arg Phe Ser Ser Leu Gln His Thr
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RAW SEQUENCE LISTING DATE: 02/19/2002
PATENT APPLICATION: US/09/925,055B TIME: 12:20:18

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Output Set: N:\CRF3\02192002\I925055B.raw

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63  Thr Leu Lys Pro Pro Asp Val Thr Cys Ile Ser Lys Val Arg Ser Ile
64  120                 125                 130                 135
66  cag atg att gtt cat cct acc ccc acg cca atc cgt gca ggc gat ggc      486
67  Gln Met Ile Val His Pro Thr Pro Thr Pro Ile Arg Ala Gly Asp Gly
68                  140                 145                 150
70  cac cgg cta acc ctg gaa gac atc ttc cat gac ctg ttc tac cac tta      534
71  His Arg Leu Thr Leu Glu Asp Ile Phe His Asp Leu Phe Tyr His Leu
72              155                 160                 165
74  gag ctc cag gtc aac cgc acc tac caa atg cac ctt gga ggg aag cag      582
75  Glu Leu Gln Val Asn Arg Thr Tyr Gln Met His Leu Gly Gly Lys Gln
76          170                 175                 180
78  aga gaa tat gag ttc ttc ggc ctg acc cct gac aca gag ttc ctt ggc      630
79  Arg Glu Tyr Glu Phe Phe Gly Leu Thr Pro Asp Thr Glu Phe Leu Gly
80      185                 190                 195
82  acc atc atg att tgc gtt ccc acc tgg gcc aag gag agt gcc ccc tac      678
83  Thr Ile Met Ile Cys Val Pro Thr Trp Ala Lys Glu Ser Ala Pro Tyr
84  200                 205                 210                 215
86  atg tgc cga gtg aag aca ctg cca gac cgg aca tgg acc tac tcc ttc      726
87  Met Cys Arg Val Lys Thr Leu Pro Asp Arg Thr Trp Thr Tyr Ser Phe
88                  220                 225                 230
90  tcc gga gcc ttc ctg ttc tcc atg ggc ttc ctc gtc gca gta ctc tgc      774
91  Ser Gly Ala Phe Leu Phe Ser Met Gly Phe Leu Val Ala Val Leu Cys
92              235                 240                 245
94  tac ctg agc tac aga tat gtc acc aag ccg cct gca cct ccc aac tcc      822
95  Tyr Leu Ser Tyr Arg Tyr Val Thr Lys Pro Pro Ala Pro Pro Asn Ser
96          250                 255                 260
98  ctg aac gtc cag cga gtc ctg act ttc cag ccg ctg cgc ttc atc cag      870
99  Leu Asn Val Gln Arg Val Leu Thr Phe Gln Pro Leu Arg Phe Ile Gln
100     265                 270                 275
102 gag cac gtc ctg atc cct gtc ttt gac ctc agc ggc ccc agc agt ctg      918
103 Glu His Val Leu Ile Pro Val Phe Asp Leu Ser Gly Pro Ser Ser Leu
104 280                 285                 290                 295
106 gcc cag cct gtc cag tac tcc cag atc agg gtg tct gga ccc agg gag      966
107 Ala Gln Pro Val Gln Tyr Ser Gln Ile Arg Val Ser Gly Pro Arg Glu
108                 300                 305                 310
110 ccc gca gga gct cca cag cgg cat agc ctg tcc gag atc acc tac tta     1014
111 Pro Ala Gly Ala Pro Gln Arg His Ser Leu Ser Glu Ile Thr Tyr Leu
112             315                 320                 325
114 ggg cag cca gac atc tcc atc ctc cag ccc tcc aac gtg cca cct ccc     1062
115 Gly Gln Pro Asp Ile Ser Ile Leu Gln Pro Ser Asn Val Pro Pro Pro
116         330                 335                 340
118 cag atc ctc tcc cca ctg tcc tat gcc cca aac gct gcc cct gag gtc     1110
119 Gln Ile Leu Ser Pro Leu Ser Tyr Ala Pro Asn Ala Ala Pro Glu Val
120     345                 350                 355
122 ggg ccc cca tcc tat gca cct cag gtg acc ccc gaa gct caa ttc cca     1158
123 Gly Pro Pro Ser Tyr Ala Pro Gln Val Thr Pro Glu Ala Gln Phe Pro
124 360                 365                 370                 375
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RAW SEQUENCE LISTING DATE: 02/19/2002
PATENT APPLICATION: US/09/925,055B TIME: 12:20:18

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Output Set: N:\CRF3\02192002\I925055B.raw

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127  Phe Tyr Ala Pro Gln Ala Ile Ser Lys Val Gln Pro Ser Ser Tyr Ala
128                  380                 385                 390
130  cct caa gcc act ccg gac agc tgg cct ccc tcc tat ggg gta tgc atg     1254
131  Pro Gln Ala Thr Pro Asp Ser Trp Pro Pro Ser Tyr Gly Val Cys Met
132              395                 400                 405
134  gaa ggt tct ggc aaa gac tcc ccc act ggg aca ctt tct agt cct aaa     1302
135  Glu Gly Ser Gly Lys Asp Ser Pro Thr Gly Thr Leu Ser Ser Pro Lys
136          410                 415                 420
138  cac ctt agg cct aaa ggt cag ctt cag aaa gag cca cca gct gga agc     1350
139  His Leu Arg Pro Lys Gly Gln Leu Gln Lys Glu Pro Pro Ala Gly Ser
140      425                 430                 435
142  tgc atg tta ggt ggc ctt tct ctg cag gag gtg acc tcc ttg gct atg     1398
143  Cys Met Leu Gly Gly Leu Ser Leu Gln Glu Val Thr Ser Leu Ala Met
144  440                 445                 450                 455
146  gag gaa tcc caa gaa gca aaa tca ttg cac cag ccc ctg ggg att tgc     1446
147  Glu Glu Ser Gln Glu Ala Lys Ser Leu His Gln Pro Leu Gly Ile Cys
148                  460                 465                 470
150  aca gac aga aca tct gac cca aat gtg cta cac agt ggg gag gaa ggg     1494
151  Thr Asp Arg Thr Ser Asp Pro Asn Val Leu His Ser Gly Glu Glu Gly
152              475                 480                 485
154  aca cca cag tac cta aag ggc cag ctc ccc ctc ctc tcc tca gtc cag     1542
155  Thr Pro Gln Tyr Leu Lys Gly Gln Leu Pro Leu Leu Ser Ser Val Gln
156          490                 495                 500
158  atc gag ggc cac ccc atg tcc ctc cct ttg caa cct cct tcc ggt cca     1590
159  Ile Glu Gly His Pro Met Ser Leu Pro Leu Gln Pro Pro Ser Gly Pro
160      505                 510                 515
162  tgt tcc ccc tcg gac caa ggt cca agt ccc tgg ggc ctg ctg gag tcc     1638
163  Cys Ser Pro Ser Asp Gln Gly Pro Ser Pro Trp Gly Leu Leu Glu Ser
164  520                 525                 530                 535
166  ctt gtg tgt ccc aag gat gaa gcc aag agc cca gcc cct gag acc tca     1686
167  Leu Val Cys Pro Lys Asp Glu Ala Lys Ser Pro Ala Pro Glu Thr Ser
168                  540                 545                 550
170  gac ctg gag cag ccc aca gaa ctg gat tct ctt ttc aga ggc ctg gcc     1734
173  Asp Leu Glu Gln Pro Thr Glu Leu Asp Ser Leu Phe Arg Gly Leu Ala
174              555                 560                 565
176  ctg act gtg cag tgg gag tcc tgaggggaat gggaaaggct tggtgcttcc       1785
177  Leu Thr Val Gln Trp Glu Ser
178          570
180  tccctgtccc tacccagtgt cacatccttg gctgtcaatc ccatgcctgc ccatgccaca   1845
181  cactctgcga tctggcctca gacgggtgcc cttgagagaa gcagagggag tggcatgcag   1905
182  ggcccctgcc atgggtgcgc tcctcaccgg aacaaagcag catgataagg actgcagcgg   1965
183  gggagctctg gggagcagct tgtgtagaca agcgcgtgct cgctgagccc tgcaaggcag   2025
184  aaatgacagt gcaaggagga aatgcaggga aactcccgag gtccagagcc ccacctccta   2085
185  acaccatgga ttcaaagtgc tcagggaatt tgcctctcct tgccccattc ctggccagtt   2145
186  tcacaatcta gctcgacaga gcatgaggcc cctgcctctt ctgtcattgt tcaaaggtgg   2205
187  gaagagagcc tggaaaagaa ccaggcctgg aaaagaacca gaaggaggct gggcagaacc   2265
188  agaacaacct gcacttctgc caaggccagg gccagcagga cggcaggact ctagggaggg   2325
189  gtgtggcctg cagctcattc ccagccaggg caactgcctg acgttgcacg atttcagctt   2385
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RAW SEQUENCE LISTING DATE: 02/19/2002
PATENT APPLICATION: US/09/925,055B TIME: 12:20:18

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192  gctgtggccc ctggacgggt acaataacac actgtactga tgtcacaact ttgcaagctc   2565
193  tgccttgggt tcagcccatc tgggctcaaa ttccagcctc accactcaca agctgtgtga   2625
194  cttcaaacaa atgaaatcag tgcccagaac ctcggtttcc tcatctgtaa tgtggggatc   2685
195  ataacaccta cctcatggag ttgtggtgaa gatgaaatga agtcatgtct ttaaagtgct   2745
196  taatagtgcc tggtacatgg gcagtgccca ataaacggta gctatttaaa aaaaaaaaaa   2805
197  aaaaaaaaaa atagcggccg cctcga                                       2831
199 <210> SEQ ID NO: 2
200 <211> LENGTH: 574
201 <212> TYPE: PRT
202 <213> ORGANISM: Homo sapien
204 <400> SEQUENCE: 2
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207  Ala Pro Glu Asp Pro Ser Asp Leu Leu Gln His Val Lys Phe Gln Ser
208              20                  25                  30
209  Ser Asn Phe Glu Asn Ile Leu Thr Trp Asp Ser Gly Pro Glu Gly Thr
210          35                  40                  45
211  Pro Asp Thr Val Tyr Ser Ile Glu Tyr Lys Thr Tyr Gly Glu Arg Asp
212      50                  55                  60
213  Trp Val Ala Lys Lys Gly Cys Gln Arg Ile Thr Arg Lys Ser Cys Asn
214  65                  70                  75                  80
215  Leu Thr Val Glu Thr Gly Asn Leu Thr Glu Leu Tyr Tyr Ala Arg Val
216                  85                  90                  95
217  Thr Ala Val Ser Ala Gly Gly Arg Ser Ala Thr Lys Met Thr Asp Arg
218              100                 105                 110
219  Phe Ser Ser Leu Gln His Thr Thr Leu Lys Pro Pro Asp Val Thr Cys
220          115                 120                 125
221  Ile Ser Lys Val Arg Ser Ile Gln Met Ile Val His Pro Thr Pro Thr
222      130                 135                 140
223  Pro Ile Arg Ala Gly Asp Gly His Arg Leu Thr Leu Glu Asp Ile Phe
224  145                 150                 155                 160
225  His Asp Leu Phe Tyr His Leu Glu Leu Gln Val Asn Arg Thr Tyr Gln
226                  165                 170                 175
227  Met His Leu Gly Gly Lys Gln Arg Glu Tyr Glu Phe Phe Gly Leu Thr
228              180                 185                 190
229  Pro Asp Thr Glu Phe Leu Gly Thr Ile Met Ile Cys Val Pro Thr Trp
230          195                 200                 205
231  Ala Lys Glu Ser Ala Pro Tyr Met Cys Arg Val Lys Thr Leu Pro Asp
232      210                 215                 220
233  Arg Thr Trp Thr Tyr Ser Phe Ser Gly Ala Phe Leu Phe Ser Met Gly
234  225                 230                 235                 240
235  Phe Leu Val Ala Val Leu Cys Tyr Leu Ser Tyr Arg Tyr Val Thr Lys
236                  245                 250                 255
237  Pro Pro Ala Pro Pro Asn Ser Leu Asn Val Gln Arg Val Leu Thr Phe
238              260                 265                 270
239  Gln Pro Leu Arg Phe Ile Gln Glu His Val Leu Ile Pro Val Phe Asp
240          275                 280                 285
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RAW SEQUENCE LISTING DATE: 02/19/2002
PATENT APPLICATION: US/09/925,055B TIME: 12:20:18

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242      290                 295                 300
243  Arg Val Ser Gly Pro Arg Glu Pro Ala Gly Ala Pro Gln Arg His Ser
244  305                 310                 315                 320
245  Leu Ser Glu Ile Thr Tyr Leu Gly Gln Pro Asp Ile Ser Ile Leu Gln
246                  325                 330                 335
247  Pro Ser Asn Val Pro Pro Pro Gln Ile Leu Ser Pro Leu Ser Tyr Ala
248              340                 345                 350
249  Pro Asn Ala Ala Pro Glu Val Gly Pro Pro Ser Tyr Ala Pro Gln Val
250          355                 360                 365
251  Thr Pro Glu Ala Gln Phe Pro Phe Tyr Ala Pro Gln Ala Ile Ser Lys
252      370                 375                 380
253  Val Gln Pro Ser Ser Tyr Ala Pro Gln Ala Thr Pro Asp Ser Trp Pro
254  385                 390                 395                 400
255  Pro Ser Tyr Gly Val Cys Met Glu Gly Ser Gly Lys Asp Ser Pro Thr
256                  405                 410                 415
259  Gly Thr Leu Ser Ser Pro Lys His Leu Arg Pro Lys Gly Gln Leu Gln
260              420                 425                 430
261  Lys Glu Pro Pro Ala Gly Ser Cys Met Leu Gly Gly Leu Ser Leu Gln
262          435                 440                 445
263  Glu Val Thr Ser Leu Ala Met Glu Glu Ser Gln Glu Ala Lys Ser Leu
264      450                 455                 460
265  His Gln Pro Leu Gly Ile Cys Thr Asp Arg Thr Ser Asp Pro Asn Val
266  465                 470                 475                 480
267  Leu His Ser Gly Glu Glu Gly Thr Pro Gln Tyr Leu Lys Gly Gln Leu
268                  485                 490                 495
269  Pro Leu Leu Ser Ser Val Gln Ile Glu Gly His Pro Met Ser Leu Pro
270              500                 505                 510
271  Leu Gln Pro Pro Ser Gly Pro Cys Ser Pro Ser Asp Gln Gly Pro Ser
272          515                 520                 525
273  Pro Trp Gly Leu Leu Glu Ser Leu Val Cys Pro Lys Asp Glu Ala Lys
274      530                 535                 540
275  Ser Pro Ala Pro Glu Thr Ser Asp Leu Glu Gln Pro Thr Glu Leu Asp
276  545                 550                 555                 560
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278                  565                 570
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281  <211> LENGTH: 211
282  <212> TYPE: PRT
283  <213> ORGANISM: Homo sapiens
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288  Asn Phe Glu Asn Ile Leu Thr Trp Asp Ser Gly Pro Glu Gly Thr Pro
289              20                  25                  30
290  Asp Thr Val Tyr Ser Ile Glu Tyr Lys Thr Tyr Gly Glu Arg Asp Trp
291          35                  40                  45
292  Val Ala Lys Lys Gly Cys Gln Arg Ile Thr Arg Lys Ser Cys Asn Leu
293      50                  55                  60
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<210> SEQ ID NO 30
<211> LENGTH: 484
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: :
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See item 11 on Ent Summary Sheet

VERIFICATION SUMMARY DATE: 02/19/2002
PATENT APPLICATION: US/09/925,055B TIME: 12:20:19

Input Set : A:\00-56 SEQ.txt
Output Set: N:\CRF3\02192002\I925055B.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1092 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1092 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: